

Seq.ST25.txt SEQUENCE LISTING

<110> Samson, Michael Parmentier, Marc Vassart, Gilbert Frederic, Libert

Frederić, Libert	
<120> Screening Methods for Identifying Compounds which Decrease HIV Entry in Cell	to a
<130> 9409/2023F	
<140> 10/661,798 <141> 2003-09-12	
<150> 09/938,703 <151> 2001-08-24	i
<150> 09/626,939 <151> 2000-07-27	
<150> 08/833,752 <151> 1997-04-09	
<150> 08/810,028 <151> 1997-03-03	
<150> EP 96870021.1 <151> 1996-03-01	
<150> EP 96870102.9 <151> 1996-08-06	
<160> 18	
<170> PatentIn version 3.1	
<210> 1 <211> 792 <212> DNA <213> Homo sapiens	
<400> 1 gaattccccc aacagagcca agctctccat ctagtggaca gggaagctag cagcaaacct 60	
tcccttcact acaaaacttc attgcttggc caaaaagaga gttaattcaa tgtagacatc 120	
tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact 180	
aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggaacaaga 240	
tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc 300	
aaaaaatcaa tgtgaagcaa atcgcagccc gcctcctgcc tccgctctac tcactggtgt 360	
tcatctttgg ttttgtgggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggc 420	
tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg tttttccttc 480	
ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt 540	
gtcaactctt gacagggctc tattttatag gcttcttctc tggaatcttc ttcatcatcc 600	

	Seq.ST25.txt
tcctgacaat cgataggtac ctggctgtcg tc	ccatgctgt gtttgcttta aaagccagga 660
cggtcacctt tggggtggtg acaagtgtga to	cacttgggt ggtggctgtg tttgcgtctc 720
tcccaggaat catctttacc agatctcaaa aa	agaaggtet teattacace tgeagetete 780
attttccata ca	792
<210> 2 <211> 1477 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1377)(1377) <223> Any nucleotide	
<220> <221> misc_feature <222> (1384)(1385) <223> Any nucleotide	
<400> 2 gaattccccc aacagagcca agctctccat ct	tagtggaca gggaagctag cagcaaacct 60
tcccttcact acaaaacttc attgcttggc ca	aaaaagaga gttaattcaa tgtagacatc 120
tatgtaggca attaaaaacc tattgatgta ta	aaaacagtt tgcattcatg gagggcaact 180
aaatacattc taggacttta taaaagatca ct	tttttattt atgcacaggg tggaacaaga 240
tggattatca agtgtcaagt ccaatctatg ac	catcaatta ttatacatcg gagccctgcc 300
aaaaaatcaa tgtgaagcaa atcgcagccc gc	cctcctgcc tccgctctac tcactggtgt 360
tcatctttgg ttttgtgggc aacatgctgg to	catcctcat cctgataaac tgcaaaaggc 420
tgaagagcat gactgacatc tacctgctca ac	cctggccat ctctgacctg tttttccttc 480
ttactgtccc cttctgggct cactatgctg co	cgcccagtg ggactttgga aatacaatgt 540
gtcaactctt gacagggctc tattttatag gc	cttcttctc tggaatcttc ttcatcatcc 600
tcctgacaat cgataggtac ctggctgtcg to	ccatgctgt gtttgcttta aaagccagga 660
cggtcacctt tggggtggtg acaagtgtga to	cacttgggt ggtggctgtg tttgcgtctc 720
tcccaggaat catctttacc agatctcaaa aa	agaaggtct tcattacacc tgcagctctc 780
attttccata cagtcagtat caattctgga ag	gaatttcca gacattaaag atagtcatct 840
tggggctggt cctgccgctg cttgtcatgg to	catctgcta ctcgggaatc ctaaaaactc 900
tgcttcggtg tcgaaatgag aagaagaggc ac	cagggctgt gaggcttatc ttcaccatca 960
tgattgttta ttttctcttc tgggctccct ac	caacattgt ccttctcctg aacaccttcc 1020
aggaattctt tggcctgaat aattgcagta go	ctctaacag gttggaccaa gctatgcagg 1080

Seq.ST25.txt	
tgacagagac tcttgggatg acgcactgct gcatcaaccc catcatctat gcctttgtcg	1140
gggagaagtt cagaaactac ctcttagtct tcttccaaaa gcacattgcc aaacgcttct	1200
gcaaatgctg ttctattttc cagcaagagg ctcccgagcg agcaagctca gtttacaccc	1260
gatccactgg ggagcaggaa atatctgtgg gcttgtgaca cggactcaag tgggctggtg	1320
acccagtcag agttgtgcac atggcttagt tttcatacac agcctgggct gggggtnggt	1380
tggnngaggt ctttttaaa aggaagttac tgttatagag ggtctaagat tcatccattt	1440
atttggcatc tgtttaaagt agattagatc cgaattc	1477
<210> 3 <211> 1442 <212> DNA <213> Homo sapiens	
<400> 3 gaattccccc aacagagcca agctctccat ctagtggaca gggaagctag cagcaaacct	60
tcccttcact acaaaacttc attgcttggc caaaaagaga gttaattcaa tgtagacatc	120
tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact	180
aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggaacaaga	240
tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc	300
aaaaaatcaa tgtgaagcaa atcgcagccc gcctcctgcc tccgctctac tcactggtgt	360
tcatctttgg ttttgtgggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggc	420
tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg ttttccttc	480
ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt	540
gtcaactctt gacagggctc tattttatag gcttcttctc tggaatcttc ttcatcatcc	600
tcctgacaat cgataggtac ctggctgtcg tccatgctgt gtttgcttta aaagccagga	660
cggtcacctt tggggtggtg acaagtgtga tcacttgggt ggtggctgtg tttgcgtctc	720
tcccaggaat catctttacc agatctcaaa aagaaggtct tcattacacc tgcagctctc	780
attttccata cattaaagat agtcatcttg gggctggtcc tgccgctgct tgtcatggtc	840
atctgctact cgggaatcct aaaaactctg cttcggtgtc gaaatgagaa gaagaggcac	900
agggctgtga ggcttatctt caccatcatg attgtttatt ttctcttctg ggctccctac	960
aacattgtcc ttctcctgaa caccttccag gaattctttg gcctgaataa ttgcagtagc	1020
tctaacaggt tggaccaagc tatgcaggtg acagagactc ttgggatgac gcactgctgc	1080
atcaacccca tcatctatgc ctttgtcggg gagaagttca gaaactacct cttagtcttc	1140
ttccaaaagc acattgccaa acgcttctgc aaatgctgtt ctattttcca gcaagaggct	1200
cccgagcgag caagctcagt ttacacccga tccactgggg agcaggaaat atctgtgggc	1260

Seq.ST25.txt ttgtgacacg gactcaagtg ggctggtgac ccagtcagag ttgtgcacat ggcttagttt
tcatacacag cctgggctgg gggtggttgg gaggtctttt ttaaaaggaa gttactgtta
tagagggtct aagattcatc catttatttg gcatctgttt aaagtagatt agatccgaat
tc
<210> 4 <211> 184 <212> PRT <213> Homo sapiens
<400> 4
Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr 1 10 15
Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu 20 25 30
Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn 35 40 45
Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met 50 60
Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu 65 70 75 80
Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe 85 90 95
Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe 100 105 110
Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu 115 120 125
Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe 130 135 140
Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser 145 150 155 160
Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr 165 170 175

Page 4

Thr Cys Ser Ser His Phe Pro Tyr 180

<210> 5 <211> 352

<211> 352 <212> PRT

<213> Homo sapiens

<400> 5

Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr 1 5 10 15

Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu 20 25 30

Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn 35 40 45

Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met 50 60

Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu 65 70 75 80

Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe 85 90 95

Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe 100 105 110

Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu 115 120 125

Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe 130 135 140

Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser 145 150 155 160

Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr 165 170 175

Thr Cys Ser Ser His Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn 180 185 190

Phe Gln Thr Leu Lys Ile Val Ile Leu Gly Leu Val Leu Pro Leu Leu 195 200 205

Val Met Val Ile Cys Tyr Ser Gly Ile Leu Lys Thr Leu Leu Arg Cys 210 220

Arg Asn Glu Lys Lys Arg His Arg Ala Val Arg Leu Ile Phe Thr Ile 225 230 235 240

Met Ile Val Tyr Phe Leu Phe Trp Ala Pro Tyr Asn Ile Val Leu Leu 245 250 255

Leu Asn Thr Phe Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser 260 265 270

Asn Arg Leu Asp Gln Ala Met Gln Val Thr Glu Thr Leu Gly Met Thr 275 280 285

His Cys Cys Ile Asn Pro Ile Ile Tyr Ala Phe Val Gly Glu Lys Phe 290 295 300

Arg Asn Tyr Leu Leu Val Phe Phe Gln Lys His Ile Ala Lys Arg Phe 305 310 315 320

Cys Lys Cys Cys Ser Ile Phe Gln Gln Glu Ala Pro Glu Arg Ala Ser 325 330 335

Ser Val Tyr Thr Arg Ser Thr Gly Glu Gln Glu Ile Ser Val Gly Leu 340 345 350

<210> 6

<211> 215

<212> PRT

<213> Homo sapiens

<400> 6

Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu 20 25 30

Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn 35 40 45

Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met 50 60

Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu 65 70 75 80

Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe 85 90 95

Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe 100 105 110

Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu 115 120 125

Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe 130 135 140

Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser 145 150 155 160

Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr 165 170 175

Thr Cys Ser Ser His Phe Pro Tyr Ile Lys Asp Ser His Leu Gly Ala 180 185 190

Gly Pro Ala Ala Ala Cys His Gly His Leu Leu Gly Asn Pro Lys 195 200 205

Asn Ser Ala Ser Val Ser Lys 210 215

<210> 7

<211> 360

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (325)..(327)

<223> Xaa = any amino acid

<400> 7

Met Leu Ser Thr Ser Arg Ser Arg Phe Ile Arg Asn Thr Asn Glu Ser $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Gly Glu Glu Val Thr Thr Phe Phe Asp Tyr Asp Tyr Gly Ala Pro Cys 20 25 30

His Lys Phe Asp Val Lys Gln Ile Gly Ala Gln Leu Leu Pro Pro Leu 35 40 45

Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn Met Leu Val Val 50 60

Leu Ile Leu Ile Asn Cys Lys Lys Leu Lys Cys Leu Thr Asp Ile Tyr Page 7 65

Leu Leu Asn Leu Ala Ile Ser Asp Leu Leu Phe Ile Ile Thr Leu Pro 85 90 95 Leu Trp Ala His Ser Ala Ala Asn Glu Trp Val Phe Gly Asn Ala Met Cys Lys Leu Phe Thr Gly Leu Tyr His Ile Gly Tyr Phe Gly Gly Ile 115 120 125 Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu Ala Ile Val His 130 140 Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe Gly Val Val 145 150 155 Ser Val Ile Thr Trp Leu Val Ala Val Phe Ala Ser Val Pro Gly Ile 165 170 175 Ile Phe Thr Lys Cys Gln Lys Glu Asp Ser Val Tyr Val Cys Gly Pro 180 185 190 Tyr Phe Pro Arg Gly Trp Asn Asn Phe His Thr Ile Met Arg Asn Ile 195 200 205 Leu Gly Leu Val Leu Pro Leu Leu Ile Met Val Ile Cys Tyr Ser Gly 210 215 220 Ile Leu Lys Thr Leu Leu Arg Cys Arg Asn Glu Lys Lys Arg His Arg 225 230 Ala Val Arg Val Ile Phe Thr Ile Met Ile Val Tyr Phe Leu Phe Trp 245 250 255 Thr Pro Tyr Asn Ile Val Ile Leu Leu Asn Thr Phe Gln Glu Phe Phe 260 265 270 Gly Leu Ser Asn Cys Glu Ser Thr Ser Gln Leu Asp Gln Ala Ile Gln 275 280 285 Val Thr Glu Thr Leu Gly Met Thr His Cys Cys Ile Asn Pro Ile Ile Tyr Ala Phe Val Gly Glu Lys Phe Arg Arg Tyr Ile Ser Val Phe 305 310 315 320

Seq.ST25.txt Arg Lys His Ile Xaa Xaa Xaa Phe Cys Lys Gln Cys Pro Val Phe Tyr Arg Glu Thr Val Asp Gly Val Thr Ser Thr Asn Thr Pro Ser Thr Gly 340 345 350 Glu Gln Glu Val Ser Ala Gly Leu <210> <211> 355 <212> PRT <213> Homo sapiens <220> <221> <222> MISC_FEATURE (231)..(233) <223> Xaa = amy amino acid <220> MISC_FEATURE <221> (333)..(335) <222> Xaa = amy amino acid <400> Met Thr Thr Ser Ile Asp Thr Val Glu Thr Phe Gly Thr Thr Ser Tyr Tyr Asp Asp Val Gly Leu Leu Cys Glu Lys Ala Asp Thr Arg Ala Leu 20 25 30 Met Ala Gln Phe Val Pro Pro Leu Tyr Ser Leu Val Phe Thr Val Gly Leu Ile Gly Asn Val Val Val Met Ile Leu Ile Lys Tyr Arg Arg 50 55 60 Ile Arg Ile Met Thr Asn Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp 65 70 75 80 Leu Leu Phe Ile Val Thr Leu Pro Phe Trp Thr His Tyr Val Arg Gly 85 90 95 His Asn Trp Val Phe Gly His Gly Met Cys Asn Leu Ile Ser Gly Phe Tyr His Thr Gly Leu Tyr Ser Glu Ile Phe Phe Ile Ile Leu Leu Thr

Seq.ST25.txt
Ile Asp Arg Tyr Leu Ala Ile Val His Ala Val Phe Ala Ile Arg Ala Arg Thr Val Thr Phe Gly Val Ile Thr Ser Ile Val Thr Trp Gly Ile 145 150 155 160 Ala Val Ile Ala Ala Leu Pro Glu Phe Ile Phe Tyr Glu Thr Glu Glu Leu Phe Glu Glu Thr Ile Cys Ser Ala Leu Tyr Pro Glu Asp Thr Val Tyr Ser Trp Arg His Phe His Thr Ile Arg Met Thr Ile Phe Cys Leu Val Leu Pro Leu Leu Val Met Ala Ile Cys Tyr Thr Gly Ile Ile Lys Thr Leu Leu Arg Cys Pro Xaa Xaa Xaa Lys Tyr Lys Ala Ile Arg Leu 225 230 235 240 Ile Phe Val Ile Met Ala Val Phe Phe Ile Glu Trp Thr Pro Tyr Asn 245 250 255 Val Ala Ile Leu Ile Ser Ser Tyr Gln Ser Leu Leu Phe Gly Asn Asn 260 265 270 Cys Glu Arg Ser Lys His Leu Asp Leu Val Met Ile Val Thr Glu Val 275 280 285 Ile Ala Tyr Ser His Cys Cys Met Asn Glu Val Ile Tyr Ala Phe Val 290 295 300 Gly Glu Arg Phe Arg Lys Tyr Ile Arg His Phe Phe His Arg His Leu 305 310 315 320 Leu Met His Leu Gly Arg Tyr Ile Pro Phe Leu Pro Xaa Xaa Ile 325 330 335 Glu Arg Ile Ser Ser Val Ser Pro Ser Thr Ala Glu Pro Glu Ile Ser

Ile Val Phe 355

<210> 9 <211> 355 <212> PRT <213> Homo sapiens

<400> 9

Met Glu Thr Pro Asn Thr Thr Glu Asp Tyr Asp Thr Thr Glu Phe 1 10 15

Asp Tyr Gly Asp Ala Thr Pro Cys Gln Lys Val Asn Glu Arg Ala Phe 20 25 30

Gly Ala Gln Leu Leu Pro Pro Leu Tyr Ser Leu Val Phe Val Ile Gly
35 40 45

Leu Val Gly Asn Ile Leu Val Val Leu Val Leu Val Gln Tyr Lys Arg 50 60

Leu Lys Asn Met Thr Ser Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp 65 70 75 80

Leu Leu Phe Ile Phe Thr Leu Pro Phe Trp Ile Asp Tyr Lys Leu Lys 85 90 95

Asp Asp Trp Val Phe Gly Asp Ala Met Cys Lys Ile Ile Ser Gly Phe 100 105 110

Tyr Tyr Thr Gly Leu Tyr Ser Glu Ile Phe Phe Ile Ile Leu Leu Thr 115 120 125

Ile Asp Arg Tyr Leu Ala Ile Val His Ala Val Phe Ala Ile Arg Ala 130 135 140

Arg Thr Val Thr Phe Gly Val Ile Thr Ser Ile Ile Ile Trp Ala Ile 145 150 155 160

Ala Ile Ile Ala Ser Met Pro Gly Leu Tyr Phe Ser Lys Thr Gln Trp 165 170 175

Glu Phe Thr His His Thr Cys Ser Leu His Phe Pro His Glu Ser Leu 180 185 190

Arg Glu Trp Lys Leu Phe Gln Ala Leu Lys Leu Asn Leu Phe Gly Leu 195 200 205

Val Leu Pro Leu Leu Val Met Ile Ile Cys Tyr Ile Gly Ile Ile Lys 210 215 220

Ile Leu Leu Arg Arg Pro Asn Glu Lys Lys Ser Lys Ala Val Arg Leu 225 230 235 240

Ile Phe Val Ile Met Ile Ile Phe Phe Leu Phe Trp Ile Pro Tyr Asn 245 250 255

Leu Thr Ile Ile Ser Val Phe Gln Asp Phe Leu Phe Thr His Glu 260 265 270

Cys Glu Gln Ser Arg His Leu Asp Leu Ala Val Gln Val Thr Glu Val 275 280 285

Ile Ala Tyr Thr His Cys Cys Val Asn Glu Val Ile Tyr Ala Phe Val 290 295 300

Gly Glu Arg Phe Arg Lys Tyr Ile Arg Gln Leu Glu His Arg Arg Val 305 310 315 320

Ala Val His Leu Val Lys Trp Leu Pro Phe Leu Ser Val Asp Arg Ile 325 330 335

Glu Arg Val Ser Ser Thr Ser Pro Ser Thr Gly Glu His Glu Ile Ser 340 345 350

Ala Gly Phe 355

<210> 10

<211> 360

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (145)..(147)

<223> Xaa = any amino acid

<220>

<221> MISC_FEATURE

<222> (321)..(323)

<223> Xaa = any amino acid

<400> 10

Met Asn Pro Thr Asp Ile Ala Asp Thr Thr Leu Asp Glu Ser Ile Tyr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Ser Asn Tyr Tyr Leu Tyr Glu Ser Ile Pro Lys Pro Cys Thr Lys Glu 20 25 30

Gly Ile Lys Ala Phe Gly Glu Leu Phe Leu Pro Pro Leu Tyr Ser Leu 35 40 45

Val Glu Val Phe Gly Leu Ile Gly Asn Ser Val Val Leu Val Leu 50 60 Phe Lys Tyr Lys Arg Ile Arg Ser Met Thr Asp Val Tyr Leu Leu Asn 65 70 75 80 Leu Ala Ile Ser Asp Leu Leu Phe Val Phe Ser Leu Pro Phe Trp Gly 85 90 95 Tyr Tyr Ala Ala Asp Gln Trp Val Phe Gly Leu Gly Ile Cys Lys Met $100 \hspace{1cm} 105 \hspace{1cm} 110$ Ile Ser Trp Met Tyr Leu Val Gly Phe Tyr Ser Gly Ile Phe Phe Val Met Ile Met Ser Ile Asp Arg Tyr Leu Ala Ile Val His Ala Val Glu 130 135 140 Xaa Xaa Xaa Ala Arg Thr Ile Ile Tyr Gly Val Ile Thr Ser Leu Ala 145 150 155 160 Thr Trp Ser Val Ala Val Phe Ala Ser Leu Pro Gly Phe Ile Phe Ser 165 170 175 Thr Cys Tyr Thr Glu Arg Asn His Thr Tyr Cys Lys Thr Lys Tyr Ser 180 185 190 Leu Asn Ser Thr Thr Trp Lys Val Leu Ser Ser Leu Glu Ile Asn Ile 195 200 205 Leu Gly Leu Val Ile Pro Leu Gly Ile Met Leu Phe Cys Tyr Ser Met 210 220 Ile Ile Arg Thr Leu Gln His Cys Lys Asn Glu Lys Lys Asn Lys Ala 225 230 235 240 Val Lys Met Ile Phe Ala Val Val Leu Phe Leu Gly Phe Trp Thr 245 250 255 Pro Tyr Asn Ile Val Leu Phe Leu Glu Thr Leu Val Glu Leu Glu Val 260 265 270 Ile Gln Asp Cys Thr Phe Glu Arg Tyr Leu Asp Tyr Ala Ile Gln Ala 275 280 285 Thr Glu Thr Leu Ala Phe Val His Cys Cys Leu Asn Pro Ile Ile Tyr 290 295 300 Page 13

Phe Phe Leu Gly Glu Lys Phe Arg Lys Tyr Ile Ile Gln Leu Phe Lys 305 310 315 320
Xaa Xaa Xaa Gly Leu Phe Val Ile Cys Gln Tyr Cys Gly Leu Leu Gln 325 330 335
Ile Tyr Ser Ala Asp Thr Pro Ser Ser Ser Tyr Thr Gln Ser Thr Met 340 345 350
Asp His Asp Leu His Asp Ala Leu 355 360
<210> 11 <211> 49 <212> PRT <213> Homo sapiens
<400> 11
Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn Phe Gln Thr Leu Lys 1 10 15
Ile Val Ile Leu Gly Leu Val Leu Pro Leu Leu Val Met Val Ile Cys 20 25 30
Tyr Ser Gly Ile Leu Lys Thr Leu Leu Arg Cys Arg Asn Glu Lys Lys 35 40 45
Arg
<210> 12 <211> 147 <212> DNA <213> Homo sapiens
<400> 12 tttccataca gtcagtatca attctggaag aatttccaga cattaaagat agtcatcttg 60
gggctggtcc tgccgctgct tgtcatggtc atctgctact cgggaatcct aaaaactctg 120
cttcggtgtc gaaatgagaa gaagagg 147
<210> 13 <211> 34 <212> PRT <213> Homo sapiens
<400> 13
Phe Pro Tyr Ile Lys Asp Ser His Leu Gly Ala Gly Pro Ala Ala Ala 1 5 10 15

Cys His Gly His Leu Leu Gly Asn Pro Lys Asn Ser Ala Ser Val 20 25 30

Ser Ly	s	
<210> <211> <212> <213>	14 27 DNA ARTIFICAL SEQUENCE	
<220> <221> <222> <223>	<pre>primer_bind (1)(27) Primer used to amplify the full size coding region of the CCR5 ne</pre>	ge
<400> tcgagg	14 atcc aagatggatt atcaagt	27
<210> <211> <212> <213>	15 27 DNA ARTIFICAL SEQUENCE	
<220> <221> <222> <223>	<pre>primer_bind (1)(27) Primer to amplify the full size coding region of the CCR5 gene</pre>	
<400> ctgatc	15 taga gccatgtgca caactct	27
<210> <211> <212> <213>	16 20 DNA ARTIFICAL SEQUENCE	
<220> <221> <222> <223>	<pre>primer_bind (1)(20) Primer used to amplify CCR5 from genomic DNA samples</pre>	
<400> cctggc	16 tgtc gtccatgctg	20
<210><211><211><212><213>	17 27 DNA ARTIFICAL SEQUENCE	
<220> <221>	primer_bind	

<222> (1)..(27)
<223> primer used to amplify CCR5 from genomic DNA samples

<400> 17 ctgatctaga gccatgtgca caactct

27

<210> 18

<211> 215

<212> PRT

<213> Homo sapiens

<400> 18

Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr 1 5 10 15

Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu 20 25 30

Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn 35 40 45

Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met 50 55 60

Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu 65 70 75 80

Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe 85 90 95

Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe 100 105 110

Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu 115 120 125

Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe 130 135 140

Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser 145 150 155 160

Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr 165 170 175

Thr Cys Ser Ser His Phe Pro Tyr Ile Lys Asp Ser His Leu Gly Ala 180 185 190 Seq.ST25.txt
Gly Pro Ala Ala Ala Cys His Gly His Leu Leu Gly Asn Pro Lys
195 200 205

Asn Ser Ala Ser Val Ser Lys 210 215